

Computation of evolutionary couplings

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 An abbreviated version of this protocol was published in eLIFE in Sep 2014

Sequence co-evolution gives 3D contacts and structures of protein complexes

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Detailed protocol

Please check the web site provided by the Marks' group: <https://evcouplings.org>

It gives access to several servers, pre-calculated data and the source code.

How to cite: (Readers should cite both the Bio-protocol preprint and the original research article where this protocol was used)

1. Bonvin, A. M. (2020). Computation of evolutionary couplings. Bio-protocol Preprint. bio-protocol.org/prep592.
2. Hopf, T. A., Schärfe, C. P. I., Rodrigues, J. P. G. L. M., Green, A. G., Kohlbacher, O., Sander, C., Bonvin, A. M. J. J. and Marks, D. S. (2014). Sequence co-evolution gives 3D contacts and structures of protein complexes. eLIFE. DOI: [10.7554/eLife.03430](https://doi.org/10.7554/eLife.03430)

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